

1 600-1-259 SEQUENCES
2
3
4 (SEQ ID No:1) LEVLFQGP
5
6
7 (SEQ ID No:2) EAIIYAAPFAKKK
8
9
10 (SEQ ID No:3) 5' primer AAA AGA AAA AAA GGC GGC CGC TCG GAT CTG
11 ATC GAA GGT CGT TGT GCG GGC AAC TTC GAC TCG G
12
13
14 (SEQ ID No:4) 3' primer GCA AAC TGG CTC TTC CGC AGC CGC TGA AGT
15 CCT CAT CGG G
16
17
18 (SEQ ID No:5) MASSRVDGGRSDLIEGRC
19
20
21 (SEQ ID No:6) CGK-[Dapa(Fl)]-GLEVLFQGPVRKG-[K^c-(Biotin)]-G-NH₂ (Cys-Fl-
22 PS-Biotin)
23
24
25 (SEQ ID No:7) PPPALPPKRRL-NH₂
26
27
28 (SEQ ID No:8)
29 Rh-KRGCAGNFDEERSSWYWGRSLRQEAVALLQQQRHGVFLVRDSSTSPGD
30 YVLSVSENSRSHYIINSSGPRPP
31 VPPSPAQPPPGVSPSRLRIGDQEFDSPALLEFYKIHYLDTTLIEPVARSQGS
32 GVILRQEEAEYVRALFDNGN
33 DEEDLPFKKGDLIRIRDKPEEQWWNAEDSEGKRGMPVPPYVEKYRPASASVS
34 ALIGGNQEGSHPQPLGGPEPGPYA
35 QPSVNTPLPNLQNGPIYARVIQKRVPNAYDKTALALEVGELVKVTKINVSGQ
36 WEGECKNGKRGHFPFTHVRLLDQQN
37 PDEDFTSGCGKDapa(Fl)GLEVLFQ
38
39
40 (SEQ ID No:9)
41 Rh-KRGCAGNFDEERSSWYWGRSLRQEAVALLQQQRHGVFLVRDSSTSPGD
42 YVLSVSENSRSHYIINSSGPRPP
43 VPPSPAQPPPGVSPSRLRIGDQEFDSPALLEFYKIHYLDTTLIEPVARSQGS
44 GVILRQEEAEYVRALFDNGN
45 DEEDLPFKKGDLIRIRDKPEEQWWNAEDSEGKRGMPVPPYVEKYRPASASVS
46 ALIGGNQEGSHPQPLGGPEPGPYA
47 QPSVNTPLPNLQNGPIYARVIQKRVPNAYDKTALALEVGELVKVTKINVSGQ
48 WEGECKNGKRGHFPFTHVRLLDQQN
49 PDEDFTSGCGKDapa(Fl)GLEVLFQGPVRKGK(Biotin)G
50